WO 2005/003156

1/35

Polypeptide			Amino acid	d sequence			SEQ ID NO:
	VDNKFNKEQQ NA	NAFYELLHLP	NINEEQRNAF	I Q SL K DDPSQ	SANLLAEAKK	LNDAQAPK	П
2222 2023 200	VDNKFNKE LR VDNKFNKE PK VDNKFNKE PR VDNKFNKE WV	QAYWEIQALP TAYWEIVKLP EAYWEIQRLP QAGSEIYNLP	nln wt o sr af nln pe orraf nln nkoka af nln raomr af	IRSLYDDPSQ IRSLYDDPSQ IRSLYDDPSQ IRSLYDDPSQ	SANLLAEAKK SANLLAEAKK SANLLAEAKK SANLLAEAKK	LNDAQAPK LNDAQAPK LNDAQAPK LNDAQAPK	ପ ଧ ୟ ପ
101 102 103 103 104 105 106 1109 1113 1114 1118 1118 1122 1123 1123 1124 1125 1125	VDNKENKEMR VDNKENKEMR VDNKENKEER	HAYWEIVKLP KAYWEIVLLP HAYWEIATLP TAYWEIVLLP TAYWEIVLLP TAYWEIVLLP TAYWEIVLLP TAYWEIVLLP TAYWEIVLLP TAYWEIVLLP TAYWEIVLLP TAYWEIVLLP NAYWEIVLLP NAYWEIVLLP TAYWEIVLLP	NLN PRQKRAF NLN NRQSRAF NLN NVQKRAF NLN PQQIRAF NLN PQQKRAF NLN NQQKRAF NLN NRQQKRAF NLN NRQ IRAF	IRSLYDDPSQ	SANLLAEAKK SANLLAEAKK	LNDAQAPK	6 10 11 11 13 14 16 17 18 19 20 21 22 23 24 25 27 28 29 29 20 21 22 23 24 27 28 28

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SEQ ID NO:	30	31	32	33	34	32	36	37	38	39	40	41			44	45	46	47			50	51				52	56	57		59	09	FIGURE 1
Amino acid sequence	NAYWEIVKLP NINPWOHRAF IRSLYDDPSQ SANLLAEAKK INDAQAPK	TAYWEIVKLP NINVROSRAF IRSLYDDPSQ SANLLAEAKK INDAQAPK	• •	NINKLOSRAF IRSLYDDPSQ SANLLAEAKK		OAYWEIVRLP NINADOKRAF IRSLYDDPSQ SANLLAEAKK INDAQAPK	NAYWEIVTLP NINKTOSRAF IRSLYDDPSQ SANLLAEAKK LNDAQAPK	NEN PGOSRAF	NLNNWOKRAF IRSLYDDPSQ SANLLAEAKK	NINKWOSRAF IRSLYDDPSQ SANLLAEAKK	NINRWOIRAF IRSLYDDPSQ SANLLAEAKK	NEN RWOTR AF	NENREONRAF	NLN NOOKR AF	NINVNOTRAF]	NLNAGOHRAF J	NEN PSOHR AF		NEN TSOTR AF	NLN VROKR AF	NENREOKRAF IRSLYDDPSQ SANLLAEAKK	NAYWEIVGLP NINQGOKRAF IRSLYDDPSQ SANLLAEAKK LNDAQAPK	NINNSORRAF IRSLYDDPSQ SANLLAEAKK	NLNSAQKRAF IRSLYDDPSQ SANLLAEAKK	NINRWOSRAF IRSLYDDPSQ SANLLAEAKK	•	EIALLP NINVAOKRAF IRSLYDDPSQ SANLLAEAKK	EIVKLP	EIVKLP	EIVSLP NINRNOSRAF IRSLYDDPSQ SANLLAEAKK	EIVKLP	
	VONKFNKELR	VDNKENKEFR	VDNKFNKENR	VDNKFNKEFR	VDNKFNKEIR	VDNKFNKEMR	VONKFNKEMR	VONKFNKEMR	VDNKFNKEMR	VDNKFNKEFR	VDNKFNKEMR	VIDNKFNKFMR	VONKFNKFLR	VONKFNKEMR	VINKFNKFLR	VDNKFNKFFR	VONKFNKFLR	VONKFNKEMR	VONKFNKELR	VONKFNKELR	VDNKFNKEPR	VINKFNKEMR	VDNKFNKEPR	VDNKENKENR	VDNKENKEMR	VONKFNKFMR	VDNKFNKEMR					
Polvpeptide	7	HERZ: 129	-HEKZ:130 7:::::0 133	ກໍເ	HERZ: 135		ZHEKZ:138	HER2:139	HER2:140	HER2:148	HER2:153	HERZ:158	HER2:159	HER2:162	HER2:165	HER2:167	HER2:168	ZHERZ:169	HER2:170	HER2:1/1	HER2:172	HER2:177	HER2:1/8	HEKZ: 1/9	-HEKZ:180 7:::530 005	HERZ: 205	HER2:207	HERZ: 209	HER2:222	HERZ: 223	ZHER2: 225	HER Z

NO:								3/	/3	5										
SEQ ID	19	62	63	64	65	99	<i>L</i> 9	89	69	70	71	72	73	74	75	16		77	78	79
Amino acid sequence	VDNKFNKE MR QAYW EI AL LP NIN RW Q IR AF I R SL Y DDPSQ SANLLAEAKK LNDAQAPK	VDNKFNKE fr tayw el av lp nln nqokr af i r sl y ddpsq sanllaeakk lndaqapk	VDNKFNKE CR TAYW EI VK LP NLN NA Q KR AF I R SL Y DDPSQ SANLLAEAKK LNDAQAPK	VDNKFNKE PK TAYW EI VV LP NLN SKQKR AF I R SL Y DDPSQ SANLLAEAKK LNDAQAPK	VDNKENKE MR NAYW EI VT LP NLN KW QIRAF IRSLYDDPSQ SANLLAEAKK LNDAQAPK	VDNKFNKE MR KAYW EI AT LP NLN KS OSRAF I R SL Y DDPSQ SANLLAEAKK LNDAQAPK	VDNKFNKE fr tayw ei vt lp nln vg q tr af i r sl y ddpsq sanllaeakk lndaqapk	KELR TAYWEIVGLP NLNTROSRAF IRSLYDDPSQ SANLLAEAKK LNDAQAPK	JDNKFNKE lr hayw ei vo lp nin re o gr af i r sl y ddpsq sanllaeakk indaqapk	VDNKFNKE FR HAYW EIIKLP NLN GKQHR AF I R SL Y DDPSQ SANLLAEAKK LNDAQAPK	KEMR TAYWEIVSLP NENTLOSRAF IRSLYDDPSQ SANLLAEAKK ENDAQAPK	/DNKFNKE MR KAYW EI QG LP NLN NR Q KR AF I R SL Y DDPSQ SANLLAEAKK LNDAQAPK	VDNKENKE MR NAYW EI AK LP NIN RE OKRAF IRSLYDDPSQ SANLLAEAKK INDAQAPK	/DNKFNKEMR HAYWEIVGLP NLNMIQORAF IRSLYDDPSQ SANLLAEAKK LNDAQAPK	IDNKENKE lr nayw ei vk lp nin ra onraf irsl y ddpsq sanllaeakk indaqapk	VDNKENKE lr tayw el ik lp nin nyorr af i r sl y ddpsq sanllaeakk indaqapk	ł	VDNKFNKE PR EAYW EI QR LP NLN NK Q KT AF I R SL Y DDPSQ SANLLAEAKK LNDAQAPK	VDNKFNKEMY AAYWEIIDLP NLNTPQIHAF IRSLYDDPSQ SANLLAEAKK LNDAQAPK	VDNKFNKE tr sayw ei vn lp nln qg qrhaf i k sl y ddpsq sanllaeakk lndaqapk
	VDNKEN	VDNKFN	VDNKFN	VDNKFN	VDNKFN	VDNKFN	VDNKFN	VDNKFNKELR	VDNKFN	VDNKEN	VDNKFNKEMR	VDNKFN	VDNKEN	VDNKEN	VDNKEN	VDNKEN		VDNKEN	VDNKFN	VDNKEN]
Polypeptide	Zuepo. 227	Zurdo220	ZHFP2.230	ZHFR2.230	Zez:252	ZHFP2.233	ZHFD2.234	ZHRP2-236	Zurdo.237	Zurdo.230	Zuepo.240	Zuedo.240	Zuebo: 264	Zueno: 204	Zueno. 271	Zuence 271	_ HERZ: 210	Zurbo.0004*	ZHER2.0434	ZHER2:3053

TGURE 1

FIGURE 2

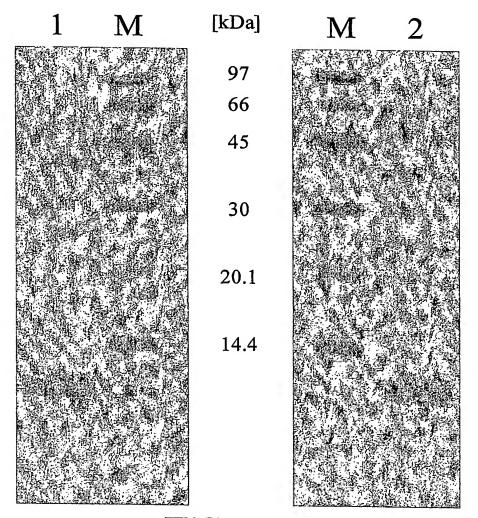


FIGURE 3

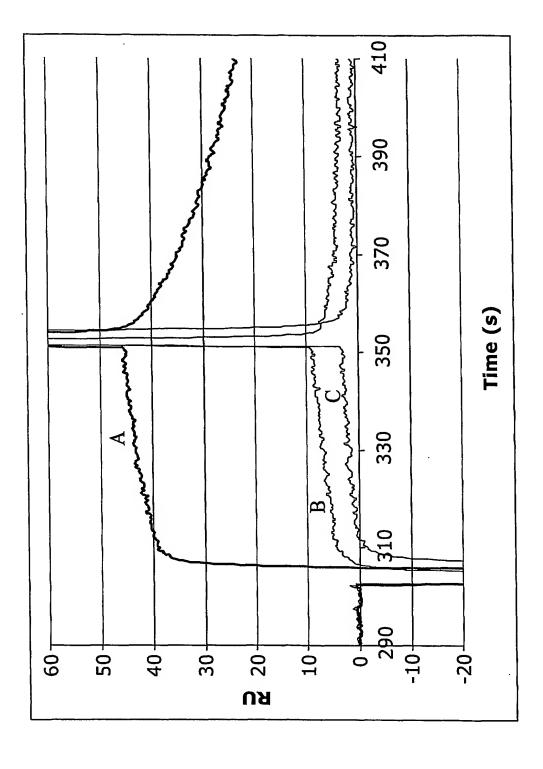
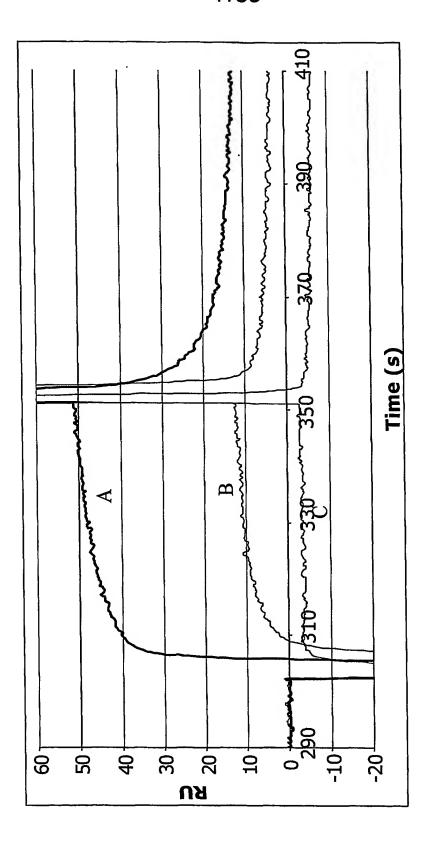


FIGURE 4

7/35

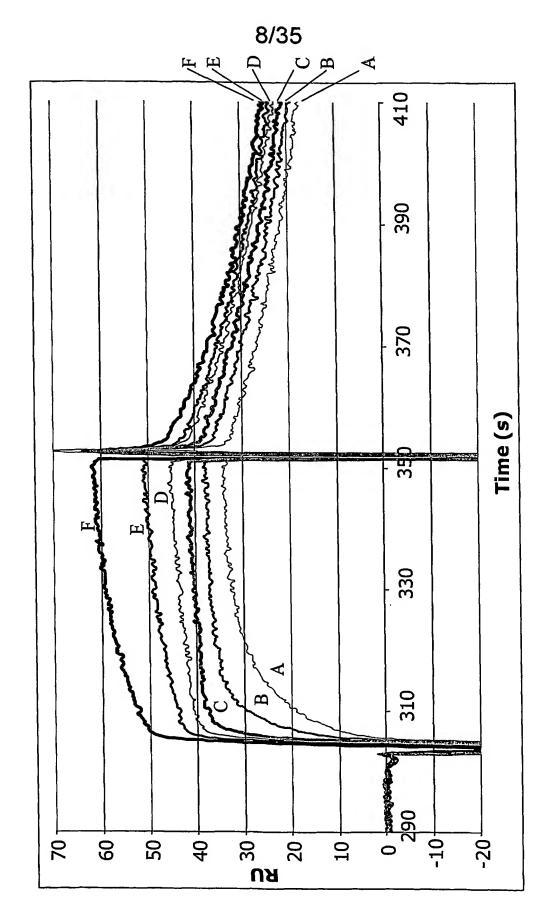


FIGIRE 5

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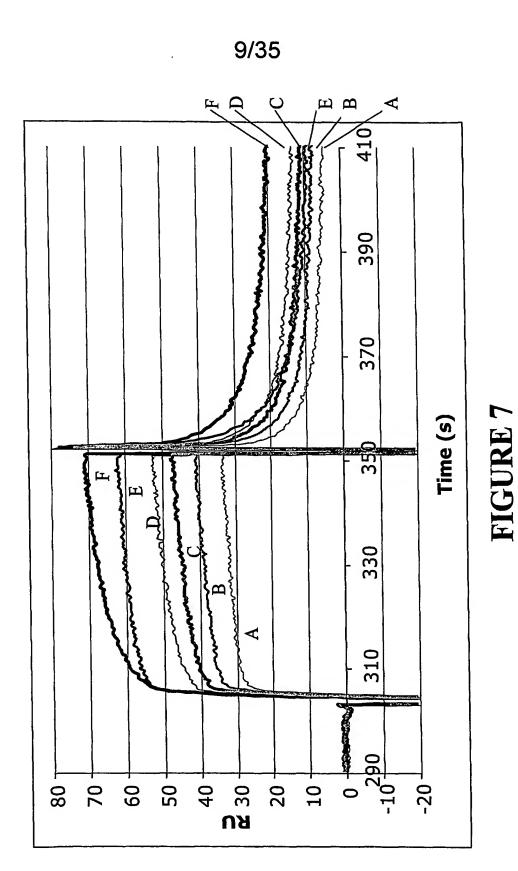
WO 2005/003156





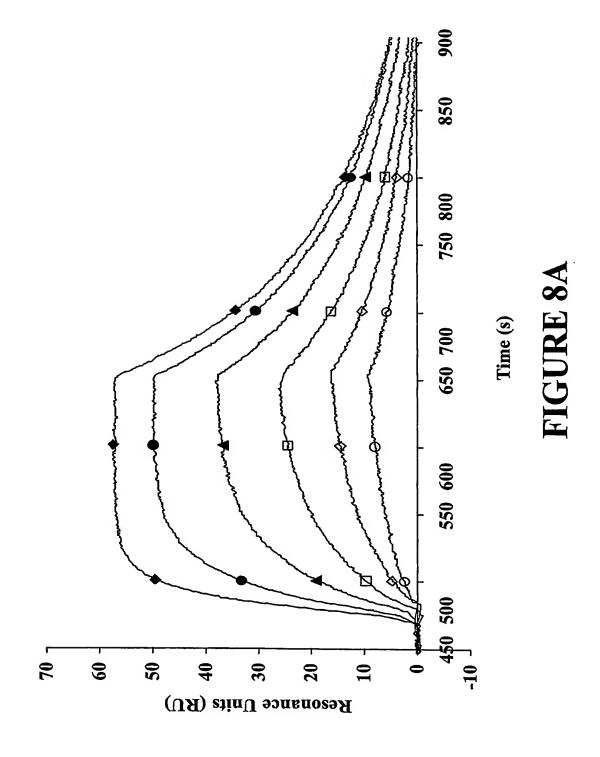
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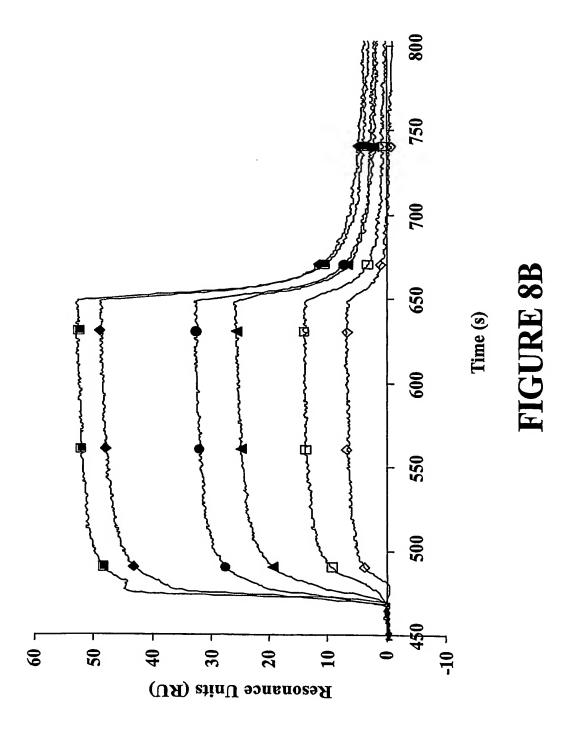


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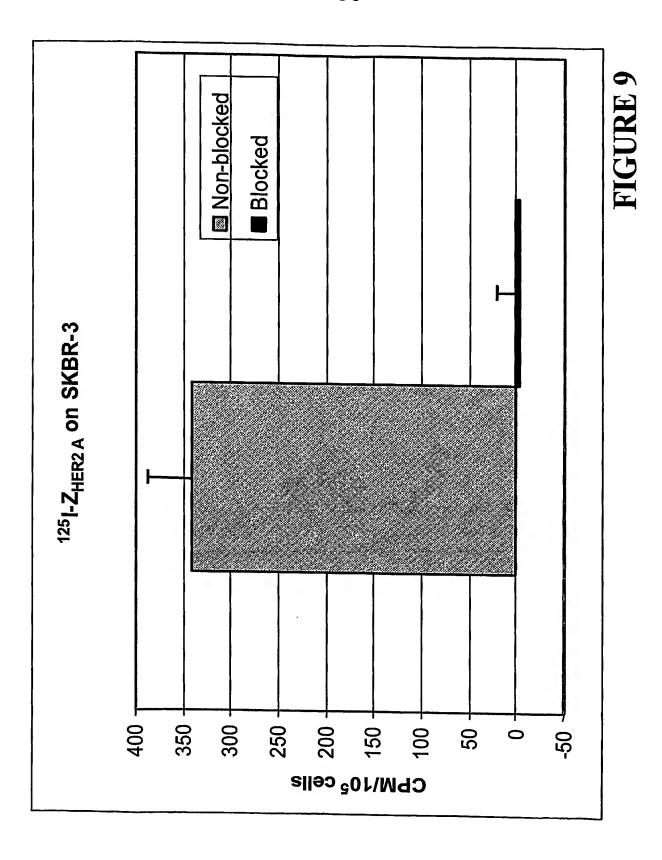
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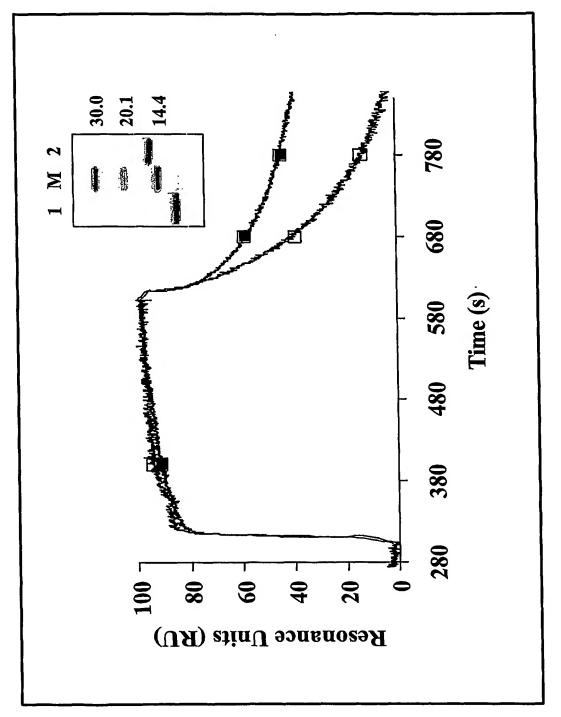


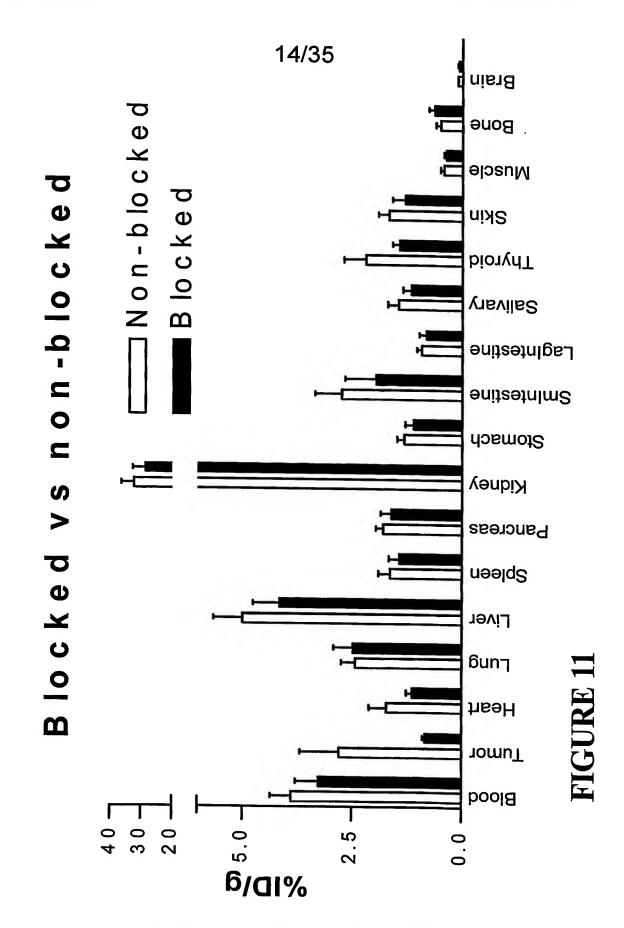
12/35



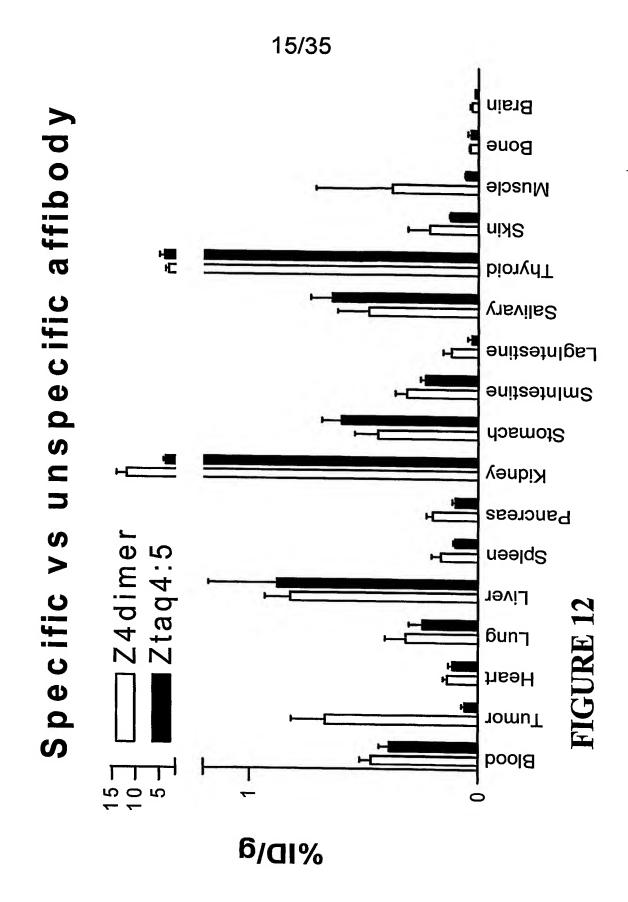
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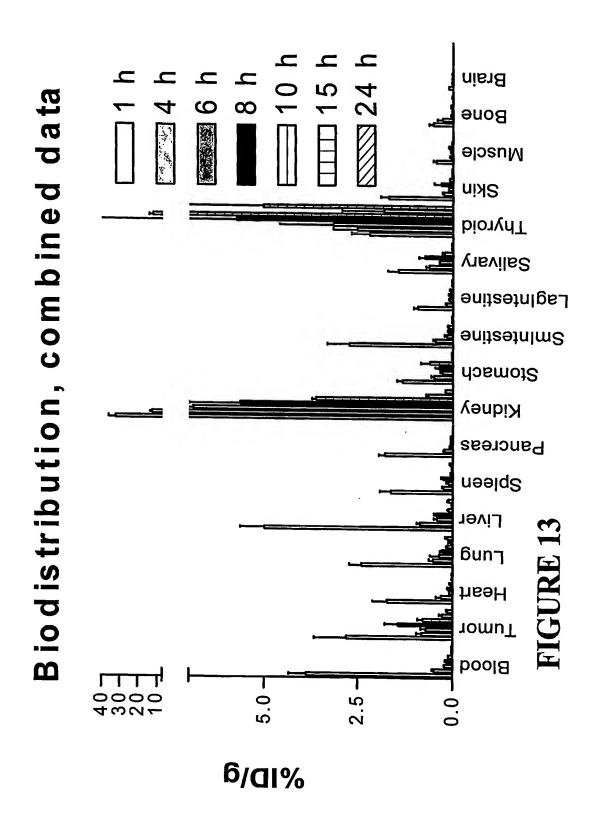




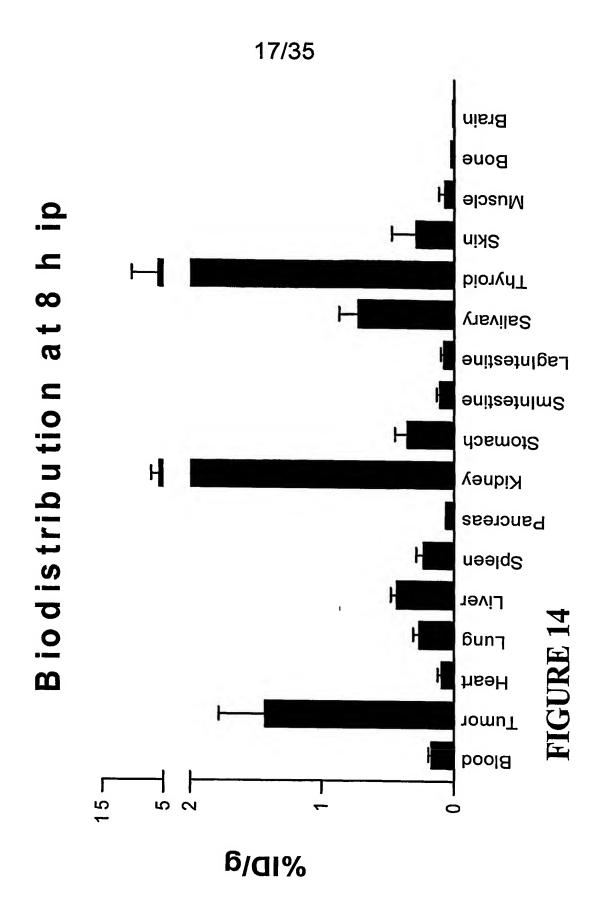
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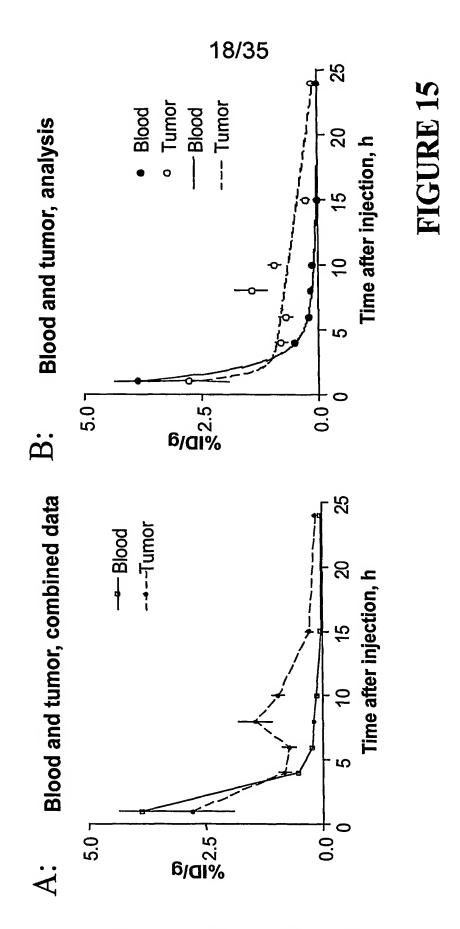
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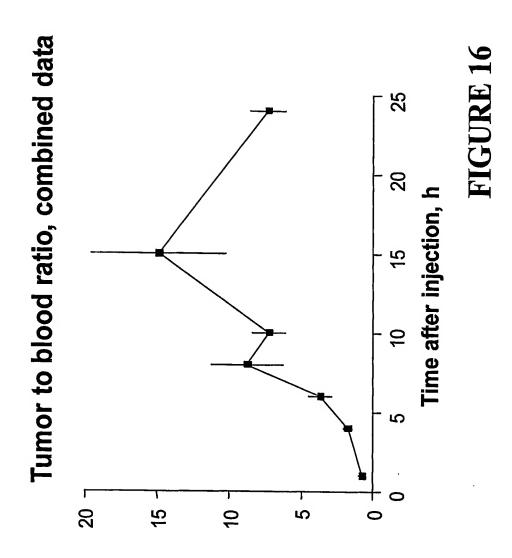
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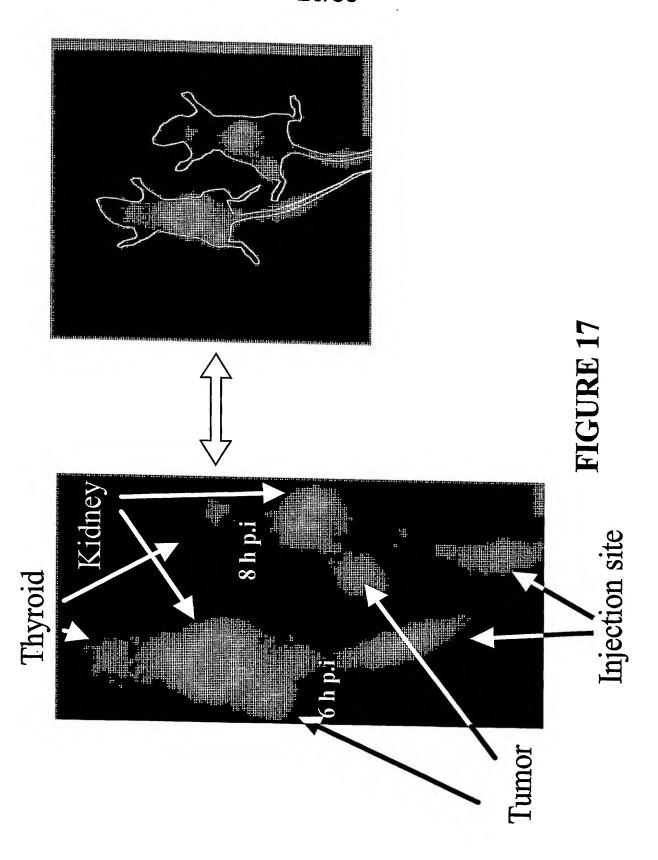
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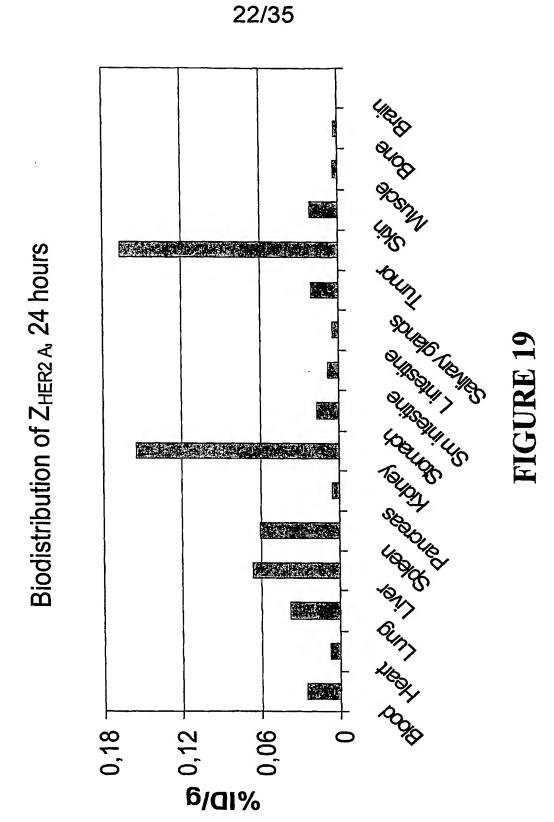


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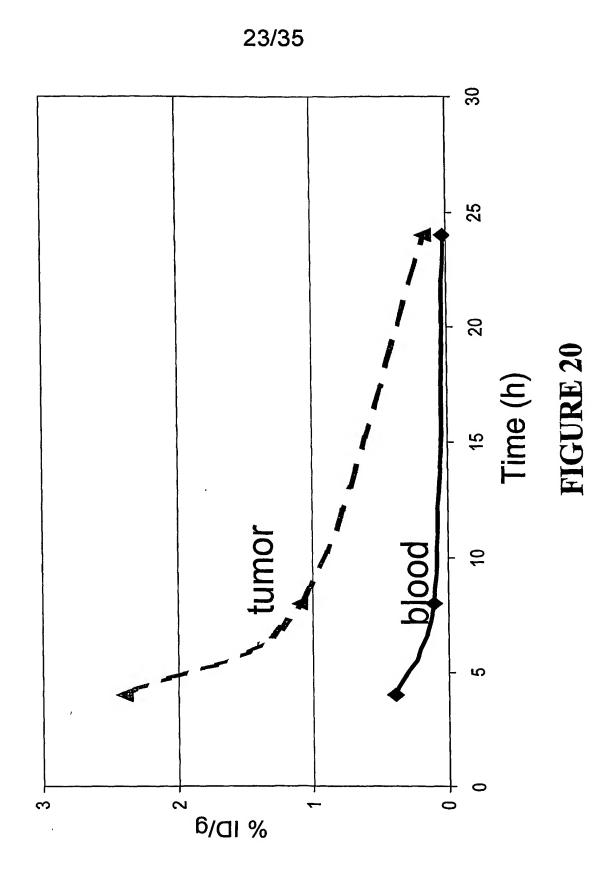
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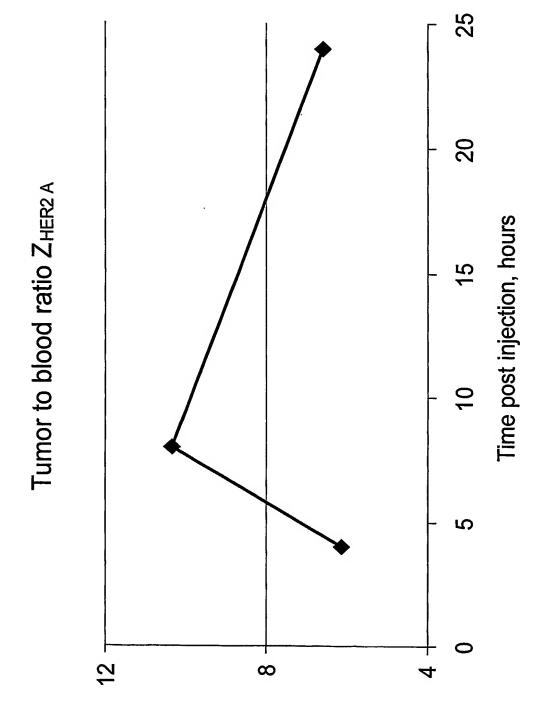
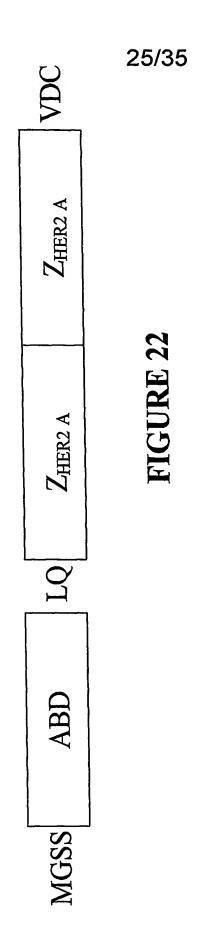
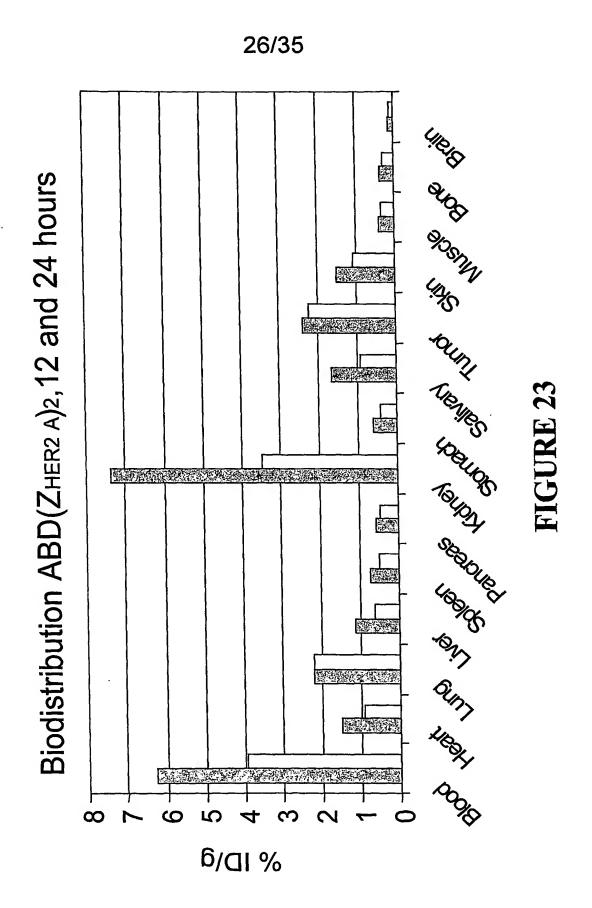


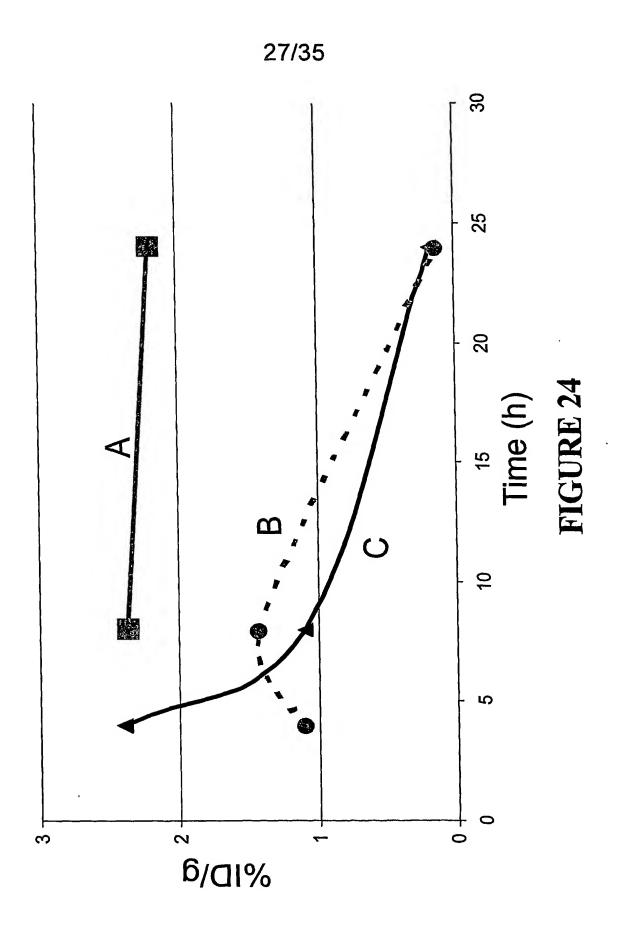
FIGURE 21



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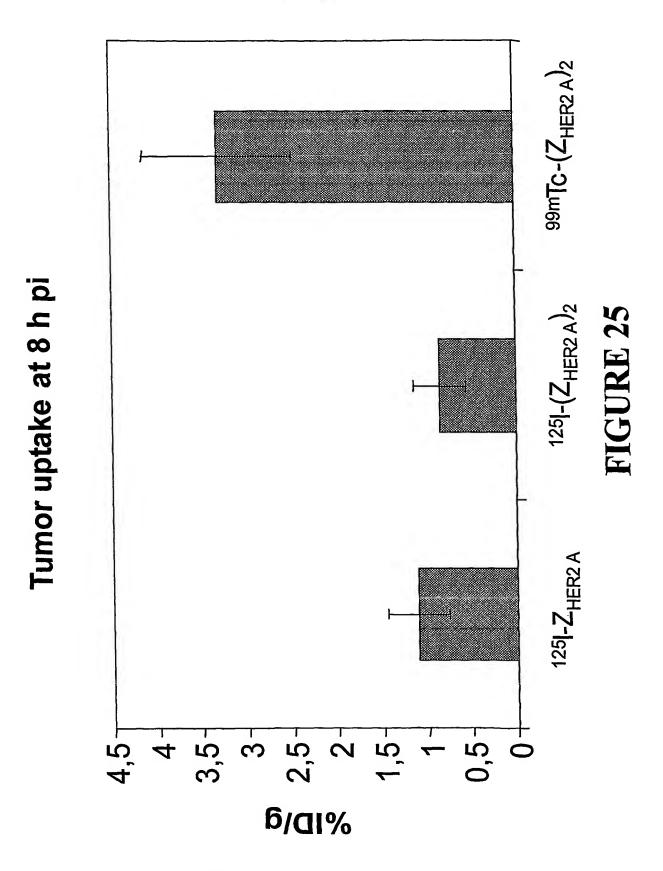


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28/35



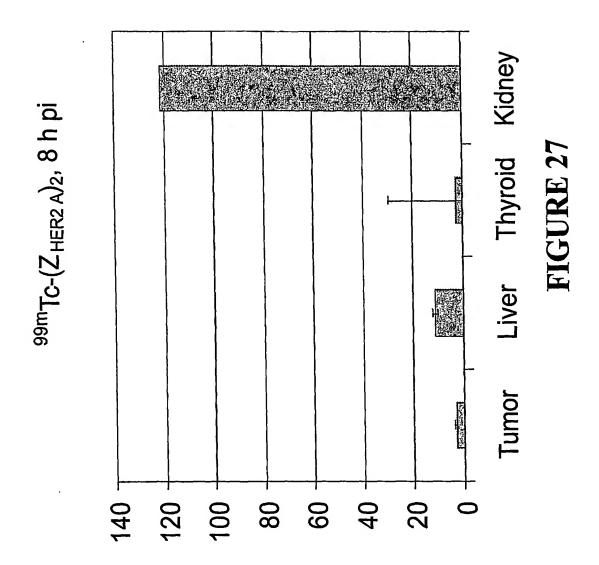
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29/35

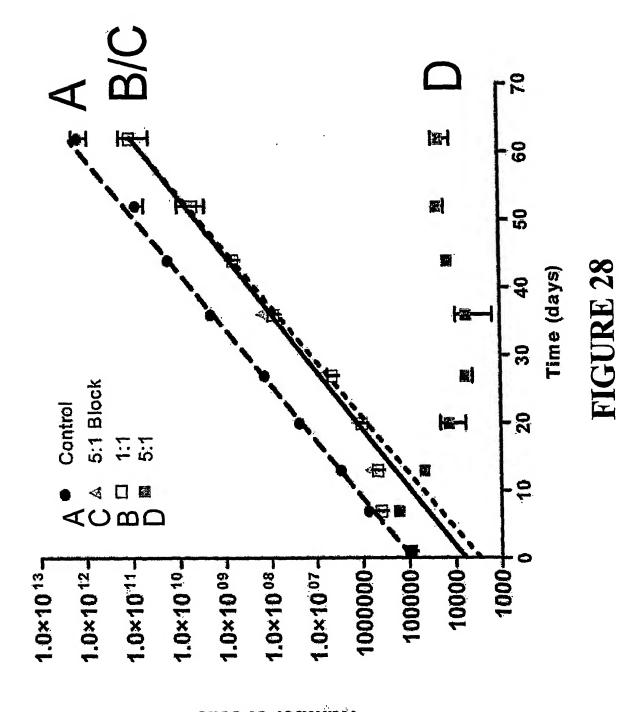
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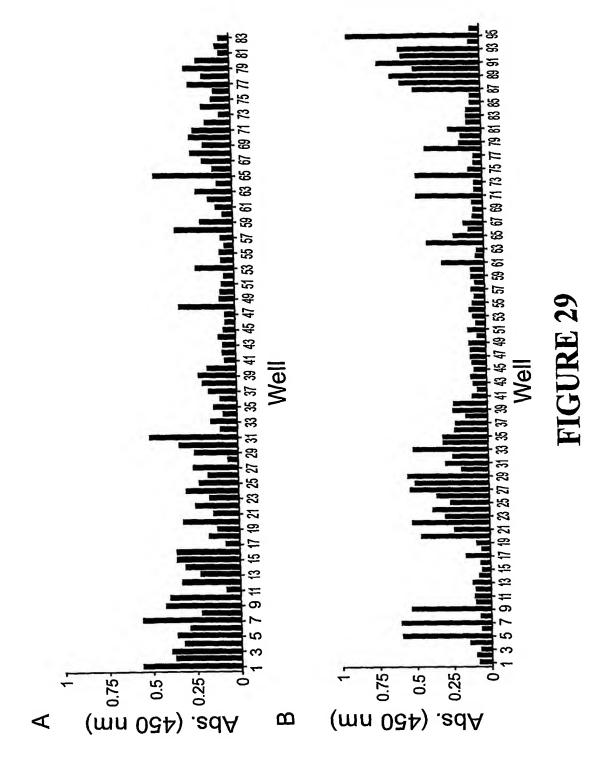
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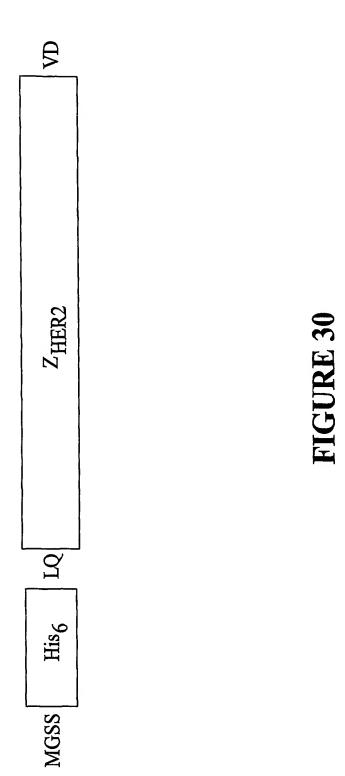


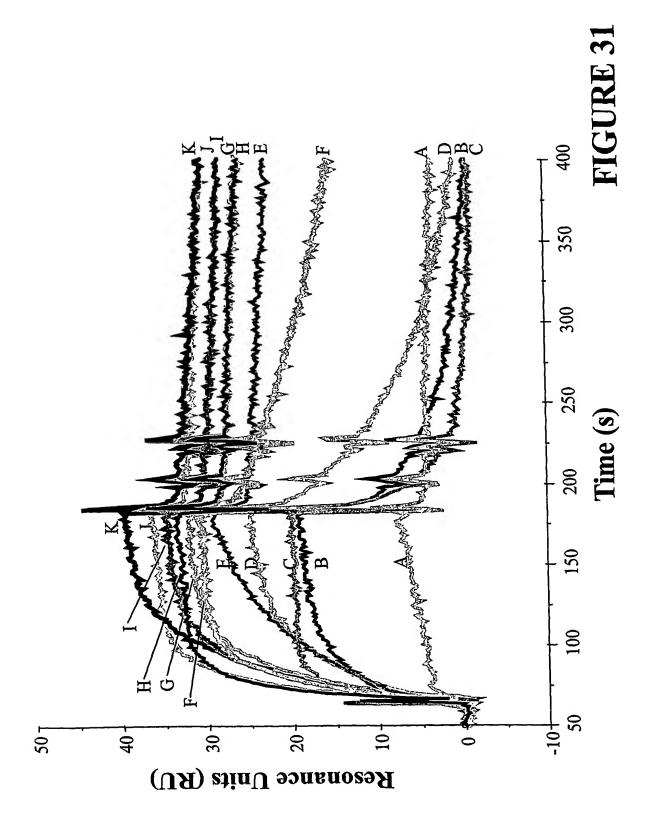
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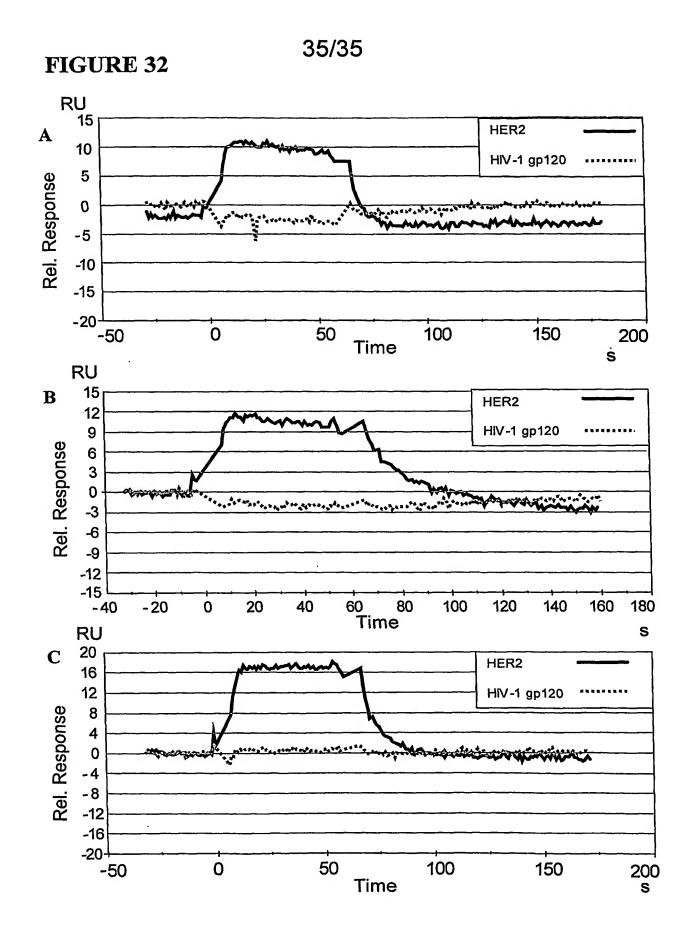
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